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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being put and is derived by analysis of the total score distribution.
   375
353
339
116.5
116.5
78.5
78.5
71.5
                                                                                                                                                                                                                            Score
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117::
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19.1
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1 /SIDS1/gcgdata/geneseq/geneseqp/AA1980.DAT:*
2: /SIDS1/gcgdata/geneseq/geneseqp/AA1981.DAT:*
3: /SIDS1/gcgdata/geneseq/geneseqp/AA1982.DAT:*
4: /SIDS1/gcgdata/geneseq/geneseqp/AA1983.DAT:*
5: /SIDS1/gcgdata/geneseq/geneseqp/AA1983.DAT:*
6: /SIDS1/gcgdata/geneseq/geneseqp/AA1985.DAT:*
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                          Length
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 625
666
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W62829
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Macadamia integrif
Macadamia integrif
Gossypium hirsutum
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Mouse liver cancer
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18.9 919 21 Y	Human androgen rec
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8.5 18.3 1162 21	HHV8 ORF 73 protei
68 18.1 326 20	urgdorferi
68 18.1 347 20	burgdorfer
18.0 489 20	G. max truncated S
67 17.9 2476 20	ig p105 zona
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5.5 17.5 1382 18	Human metastasis-a
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63 16.8 371 20	
3 16.8 1251 16	Mouse LTBP-3. Mus
2.5 16.7 436 17	Human 70K U1 snRNP
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2.5 16.7 594 1/ 2.5 16.7 595 17	SAP-ALAMET-VEGF165
5 16.7 614 16	70K autoantigen, p
2.5 16.7 671 21	Human PRO1604 (UNO
1 16.3 71 20	e seq II
1 16.3 86 20	GST-HD fusion prot
1 16.3 499 9 1	storage pi
61 16.3 910 20 Y22 0 5 16 1 2482 16 R72	brain CNG
5 16.1 2482 19 W2399	Human mitosin amin
0.5 16.1 3248 17 R9979	Kinetochore protei
16.0 537 16 R7518	Osteoinductive ret
0 16.0 1233 20 Y559	
ALIGNMENTS	
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52830 standard; Protein; 625 AA.	
52830;	
)-OCT-1998 (first entry)	
scadamia integrifolia antimicrobial protein.	
ntimicrobial protein; infestation; control.	
acadamia integrifolia.	
Y Location/Qualifiers	

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RESULT
W62830
                                                                                                                      Key
Peptide
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N-PSDB; V42316
      WPI; 1998-377279/32
                   Bower NI, Goulter KC,
                                                                       02-JUL-1998.
                                (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
                                              20-DEC-1996;
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                                                                                     WO9827805-A1.
                                                                                                         Protein
                                              96AU-0004275.
                                                          97WO-AU00874
                                                                                                        /note= "signal peptide" 29..666
                                                                                                 "mature protein
                   Green JL,
                   Manners JM,
                   Marcus JP;
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RESULT
W62829
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Sequence
                     animals.
                             The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and map {\bf map}
                                                              Claim 1; Page 39-41; 96pp; English
                                                                                   Novel anti-microbial protein from e.g. Macadamia useful for controlling microbial infestations of
                                                                                                                                                                                                                                           02-JUL-1998
                                                                                                                                                                                                                                                                                                 Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals
                                                                                                                     WPI; 1998-377279/32.
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666 AA;
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                                                                                                                                                                                                                                                                                   /note= "mature protein"
                                                                                                                                                                                                                                                                                                          /note= "signal peptide"
                                                                                                                                                                                                                                                                                                 .666
                                                                                                                                                    Green JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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Pred. No. 9.1e-36;
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                                                                                                                                                    Manners JM,
                                                                                                                                                     Marcus
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plants or man
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Query Match

94.18;

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RESULT

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RESULT
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ID W6:
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Best Local Similarity 89.4
Matches 59; Conservative
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N-PSDB; V42310.
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| 246 kqsdnp 251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 339; DB 19;
Pred. No. 1.5e-31;
4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 3.5e-33; 2; Mismatches 2;
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Best Local S
Matches 36
                                                                                                                                                                                                        Theobroma cacao
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20-DEC-1996;
                                                  22-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         141 kqqcvrecrekyqenpwrgereeeaeeeeteegeqeqshnp 181
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nes 36; Conserv
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Pred. No. 2.9e-07;
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plants or mammals
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The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocao beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close
                                                                                                                                                                                                                                                                                                                                         Claim 4; Fig 2; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Recombinant cocoa proteins - are responsible for flavour in cocoa beans and produced in large quantities using yeast and bacterial expression vectors
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N-PSDB; Q20377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spencer ME,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cocoa; flavour; vicilin; seed storage protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence encoded by 67 kD T. cacao protein cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MRSC ) MARS UK LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-DEC-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 47-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-377279/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               141 nnp 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     64 DNP 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                82 qrqyqqcqgrcqeqqqqqqqqqcqrkcweqykeqer-gehenyhnhkknrseeeeggqr 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 QREYEDCRRHCEQQE--PRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGSGRYEEGEEKQS 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          σ
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l Similarity 39.7%;
25; Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Goulter KC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       525 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hodge R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90GB-0013016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91WO-GB00914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12;
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Pred. No. 7.3e
12; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ΕA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ashton
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.3e-06;
nes 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marcus
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         revealed close
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Best Local S
Matches 25
                                                                                                                                      Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                           The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and \mbox{\it man}
                                                                                                                                                                                                                                                                                    Claim 1; Page 60-62; 96pp; English.
                                                                                                                                                                                                                                                                                                            Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals
                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antimicrobial protein; infestation; control.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        homologies between the 67 kD polypeptide and the vicilins, which are seed storage proteins.
                                                                                                                                                                                                                                                                                                                                                                                                     (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hordeum vulgare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-OCT-1998 (first entry)
                                 |: ||| ::
102 rhgegeree 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               141 nnp 143
                                                 54 RYEEGEEKQ 62
|: ||| ::
                                                                                       42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 31.1%; Score 116.5; DB 13; Length 566;
Local Similarity 39.7%; Pred. No. 7.9e-06;
Nes 25; Conservative 12; Mismatches 23; Indels 3;
                                                                                                               10 EDCRRHCEQQEPRLQY-QCQRRCQEQQRQHGRGGDLMNPQRG------GSG 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         64 DNP 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 QREYEDCRRHCEQQE--PRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGSGRYEEGEEKQS 63
œ
                                                                                      qqcvqrcrqerprysharcvqecrddqqqhgrheqeeeqgrgrgwhgegereeehgrgrg 101
                                                                                                                                                                                                                                                                                                                                                                           Goulter KC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  566 AA;
                                                                                                                                                                                                          637 AA;
                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               96AU-0004275
                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-AU00874
                                                                                                                                        25.3%; Score 95; DB 19; 1
29.0%; Pred. No. 0.0028;
tive 12; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                          Green JL,
                                                                                                                                                                                                                                                                                                                                                                           Manners JM,
                                                                                                                                                                 DB 19; Length 637;
                                                                                                                                                                                                                                                                                                                                                                             Marcus JP;
                                                                                                                                         Indels
                                                                                                                                                                                                                                               and mammalian
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Best Local Similarity
Matches 20; Conserv
                                                                                                           W37483 standard; Protein; 669
                                                                                                                                                                                                                                                                                                                                                                                                                               The protein is used to raise antibodies for receptor assays and for affinity purification. The 98 kD product starts at the first Met codon; the 79 kD product starts from the second.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P90996 standard; protein; 919 AA
             Mouse liver cancer-originated culture cell growth factor.
                                            27-MAR-1998 (first entry)
                                                                             W37483;
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 8; Fig 3; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           purification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1989-309501/42.
N-PSDB; N91577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liao S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAR-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO8909223-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Androgen receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human androgen receptor DNA clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-1990 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P90996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New DNA encoding new androgen receptor and TR2 polypeptide(s) - able to bind DNA, and derived antibodies, useful for receptor assay and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ARCH-) ARCH DEVELOPMENT CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-MAR-1989;
                                                                                                                                                                                       113 qp 114
                                                                                                                                                                                                                       65 NP 66
                                                                                                                                                                                                                                                     58 qqqqqqq....-qqqqqqqqqqtsptqqqqqqqqqbgedgspqahrrgptgylvldeeqqps 112
                                                                                                                                                                                                                                                                       5 QQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGSGRYEEGEEKQSD 64
                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chang C;
                                                                                                                                                                                                                                                                                                                                                                                                   919 AA;
                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88US-0176107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89WO-US01238
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/product=98 kD polypeptide
185..919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
1..919
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                                                                                                                                                                                                                                                                                                                20.9%; Score 78.5; DB 10; 32.3%; Pred. No. 0.34; ative 11; Mismatches 26;
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                                                                                                                                                                                                                                                                                                                    26;
                                                                                                                                                                                                                                                                                                                     Indels
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RESULT
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide comprising at least 5 continuous amino acids; (2) a polypeptide protein similar to protein of (1), but comprises no protein combined naturally and also has 2 amino acid sequences (as given in the specification) which may be modified to produce a sequence of at least 5 amino acids; (3) a recombinant DNA molecule comprising a vector and the DNA segment of (1); (4) a cell comprising (3); (5) producing a polypeptide comprising amino acids corresponding to a protein of (1), in which a cell comprising (3) is cultured under a conditions enabling the expression of (1) to produce the polypeptide which is isolated. The above method may be used to produce proteins HET-A and HET-B.
                                                                      Human; trichohyalin; TRHY; protein; tissue structure; wound healing; terminally differentiating epidermal tissue; proteinaceous gel;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a protein which has local homology to human liver cancer-originated culture cell isolated from a mouse testicle cDNA library. The sequence may (1) be modified to produce a polypeptide comprising at least 5 continuous amino acids; (2) a
                         Homo sapiens
                                                                                                                 A human trichohyalin (TRHY) protein.
                                                                                                                                                   25-NOV-1999
                                                                                                                                                                                  Y30795;
                                                                                                                                                                                                              Y30795 standard; Protein; 1898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA segment encoding protein homologous to human liver cancer-originated culture cell - which may be modified to polypeptide of at least 5 continuous amino acids, useful i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse;
hHDGF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 10; Page 13-14; 18pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-080076/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP09313185-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        producing, e.g. protein HET-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB; V01731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (KISH/) KISHIMOTO C.
(SEKI ) SEKISUI CHEM IND CO LTD.
                                                                                                                                                                                                                                                                                        371 ledeep 376
                                                                                                                                                                                                                                                                                                                                                    322 rrdeerrrelearrrreqeeelrrlreqer-eekerrkera----
                                                                                                                                                                                                                                             10
                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                        61 KQSDNP
                                                                                                                                                                                                                                                                                                                                                                     1 KRDPQQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGKGGDLMNPQRGGSGRYEEGEE 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human; liver cancer-originated culture cell growth factor, \mbox{HET-A}; \mbox{HET-B}.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              669 AA;
                                                                                                                                                                                                                                                                                                                        66
                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96JP-0131788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96JP-0131788
                                                                                                                                                                                                                                                                                                                                                                                                                          19.6%;
31.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                            Score 73.5; DB Pred. No. 0.92;
                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                    --erggs----sgee 370
                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 11
P93109
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Best Local Sim
Matches 20;
DNA encoding androgen receptor protein - useful for transforming eukaryotic hosts for protein expression and subsequent antibody prodn.
                                                                                                                                                                                                                                                                                                                                                              1112 reepekrrrgerergcreeeelqgeeegllreerekrrrgelergyreeeelgrgkr--k 1169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a human trichohyalin (TRHY) protein. The protein is found in terminally differentiating epidermal tissue, and is involved in forming the structural architecture of such tissue. The trichohyalin protein is useful for forming a proteinaceous gel which may then be used for healing wounds, or in breast implants.
                                                                                                                                                                                                                                                                                                           P93109 standard; protein; 919 AA
                                N-PSDB; N91772
                                           WPI; 1989-324206/44.
                                                                 French FS,
                                                                                                                                13-APR-1989;
                                                                                                                                                      19-OCT-1989
                                                                                                                                                                           WO8909791-A
                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                    Human androgen
                                                                                                                                                                                                                                          Human androgen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-561041/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kim I,
                                                                                      (UYNC-) UNIVERSITY OF NORTH CAROLINA
                                                                                                            14-APR-1988;
                                                                                                                                                                                                                                                               19-MAR-1990 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (USSH ) US DEPT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                      53 GRYEEGEEKQSD
                                                                                                                                                                                                                                                                                                                                                                                                                              1 KRDPQQREYEDCRRHCEQQEPRLQYQCQ-----RRCQEQQRQHGRGGDLMNPQRGGS 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        trichohyalin useful for forming
                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 19.1%; Score 71.5; DB Similarity 27.8%; Pred. No. 4.7; 20; Conservative 17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chung S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1898 AA;
                                                                Wilson
                                                                                                          88US-0182646
                                                                                                                                89WO-US01548
                                                                                                                                                                                                                    receptor; ployclonal antibody; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9308-0056200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9705-0800644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3A-W; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                      64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Park
                                                                 E
M
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                                                                Joseph DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Steinert PM,
                                                                 Lubahn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proteinaceous gel that promotes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                     24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     11;
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RESULT
W14783
ID W1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             망
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                     prostatic cells. Antisense oligonucleotides (see also T63200, T63404-05) based on an androgen receptor CDNA clone (see also T63407) can be used to prevent androgen receptor gene expression, thereby inhibiting the growth or survival of prostatic cells for the treatment of benign prostatic hyperplasia and prostate cancer
 Sequence
                                                                                            Human androgen receptor (W14783) binds testosterone and, acting at the transcriptional level, regulates the growth of normal
                                                                                                                                      Disclosure; Page 22-28;
                                                                                                                                                                             Oligonucleotide(s) antisense to human androgen receptor and acidic FGF genes - used to inhibit gene expression, for the treatment of
                                                                                                                                                                                                                         WPI; 1997-202879/18
N-PSDB; T63407.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Androgen receptor; acidic fibroblast growth factor; aFGF; antisense; benign prostatic hyperplasia; prostate cancer; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the presence of endogenous androgen, as androgen will not interfere with binding. They may be used in assays to determine and quantify cellular distribution of AR in tumour tissue, and are esp. useful for evaluating
                                                                                                                                                                   benign prostatic hyperplasia
                                                                                                                                                                                                                                                                                                                                                    20-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                              27-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Androgen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W14783
                                                                                                                                                                                                                                                                                             (WORC-) WORCESTER FOUND BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                        20-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W14783 standard; Protein; 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig. 4; 41pp; English.
                                                                                                                                                                                                                                                                   Zamecnik PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prostate cancers to determine responsiveness to androgen withdrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           116 qp 117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGRGGGLMNPQRGGGGRYEEGEEKQSD 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NP 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                qqqqqqqqqqqqqqqqqqqetspr--qqqqqqqedgspqahrrgptgylvldeeqqps 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor protein (AR) is used to produce mono- or poly-clonal es. These are used for the detection and quantification of AR in
 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA;
                                                                                                                                                                                                                                                                                                                        95US-0004018
                                                                                                                                                                                                                                                                                                                                                    96WO-US15081
                                                                               Antisense oligonucleotides (see a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18.9%;
                                                                                                                                     51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 71;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 919;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          not interfere with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2;
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RESULT
Y78914
ID Y7
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                                           This sequence represents the human androgen receptor (AR) amino acid sequence. The invention relates to a fragment of the AR corresponding to CC amino acids 234-391 (see Y78913). The fragment mediates are considered activation of the AR. The androgen receptor acts as CC androgen independent activation of the AR. The androgen receptor acts as CC androgen responsive genes. Interaction of the AR with the protein kinase CC androgen recepton pathway involves interaction with the androgen CC independent region. The AR fragment and peptides derived from it can be CC used as agents for inhibiting androgen independent activation of the AR. The CC androgen receptor, as activation domains, and as a tool for screening for CC compounds which affect androgen independent activation of the AR. The CC compounds which affect androgen independent activation of the AR. The CC init androgen mediated disease progression. These diseases include CC cancer, benign prostatic hyperplasia, hirusutism, androgenic alopecia, CC cancer, breast cancer, Kennedy disease, and especially prostate cancer. The peptides and nucleic acids encoding them, are especially used for the CC treatment of androgen mediated diseases, especially prostate tumours in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Вþ
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Best Local S
Matches 18
Sequence
                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 7; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                              screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel non-androgen ligand binding peptides for inhibiting androgen-independent activation of androgen receptor, use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-182113/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acne; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Androgen receptor; AR; androgen-independent activation; inhibitor; cancer; benign prostatic hyperplasia; hirusutism; androgenic alopecia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human androgen receptor (AR) amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Y78914;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Y78914 standard; protein; 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sadar MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    y Match 18.9%; Score 71; DB Local Similarity 29.0%; Pred. No. 2.5; hes 18; Conservative 14; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGGGRYEEGEEKQSD 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              qqqqqqqqqqqqqqqqqqqetspr--qqqqqqgedgspqahrrgptgylvldeeqqps 115
                                  deprived of androgen.
                                                                                                                                                                                                                                                                                                                                                                                                                                              compounds and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bruchovsky N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer; Kennedy disease; prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0091871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-CA00604
                                                                                                                                                                                                                                                                                                                                                                                                                            cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                              for treatment of androgen-mediated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Snoek R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18; Length 919;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 used
                                                                                                                                                                                                                                                                                                                                                                                                                                              diseases
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Y33504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ş
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  Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                This invention describes novel pure proapoptotic dependence peptides which comprise a sequence of an active dependence domain selected from dependence polypeptides consisting of p75NTR, androgen receptor, DCC, nuntingtin polypeptide, Machado-Joseph disease gene product, SCA1, SCA2, SCA6 and atrophin-1 polypeptide. The proapoptotic peptides are capable of inducing cell death and can be used to develop products to mediate or inhibit apoptosis. The methods can be used for reducing the severity of a proapoptotic dependence domain mediated pathological conditions e.g. Huntington's disease, Alzheimer's disease, Kennedy's disease, Schotstonburger, Sch
                                                                                                                                                                                          Spinocerebellar ataxias, dentatorubropallidoluysian atrophy, Machado-Joseph disease, stroke or head trauma. They can also be used for reducing the severity of a pathological condition mediated by upregulated cell proliferation or cell survival e.g. neoplastic, malignant, untoimmune or fibrotic conditions. This sequence represents a human unliganded androgen receptor described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Proapoptotic; dependence domain; p75NTR; androgen receptor; DCC; huntingtin polypeptide; Machado-Joseph disease; SCA1; SCA2; SCA6; atrophin-1; cell death; apoptosis; Huntington's disease; head trauma; Alzheimer's disease; Kennedy's disease; spinocerebellar ataxia; stroke; dentatorubropallidoluysian atrophy; cell proliferation; cell survival;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New proapoptotic dependence peptides, used to develop products for treating, e.g. Alzheimer's disease -
                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 178-179; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bredesen DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BURN-) BURNHAM INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9945944-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neoplastic; malignant; autoimmune;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Y33504 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human unliganded androgen receptor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           116 qp 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65 NP 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58 qqqqqqqqqqqqqqqqqqqqtspr--qqqqqqgedgspqahrrgptgylvldeeqqps 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 QQREYEDCRRHCEQQEPRLQYQCQRRCQEQQRQHGRGGDLMNPQRGGSGRYEEGEEKQSD 64
        . Similarity
19; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e.g. Alzheimer's disease
                                                                                                                                          154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                         18.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14;
Score 69.5; DB Pred. No. 0.57; 9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 71; I
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 21; Length 919; 2.5;
                                                        DB 20;
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                                                     Length 154;
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Matches 19
                                                                                                                            the DNA-binding domain it is identical to that of rAR DNA-binding domain. Homology comparisons with other known steroid receptors indicate that hAR is more closely related to glucocorticoid, mineralo-corticoid and progesterone receptors than to v-erb-A or to receptors for oestrogen, vitamin D and thyroid hormones.
                                                                                                        Sequence
                                                                                                                                                                                    This sequence was deduced from a cDNA clone isolated by screening commercially available human testis and prostate lambda gtll cDNI libraries. The sequence is very similar to that of rat AR and in
                                                                                                                                                                                                                                    Claim 25; Fig 3; 79pp; English.
                                                                                                                                                                                                                                                         Androgen receptor and TR2 DNA binding proteins - DNA sequences and antibodies for detection and quantification methods
                                                                                                                                                                                                                                                                                                                                  Liao S,
                                                                                                                                                                                                                                                                                                N-PSDB; Q12001.
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                                                                                                                                                                                                                                                                                                                                                                                                        19-OCT-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hAR; DNA-binding protein; steroid hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human androgen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R12223;
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                        17
                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 59 qqqqqqqqqqqqqqqqqqqqqqqqqqqqqhrrqptqylvldeeqqpsqp 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15
qqqqqqqqqqqqqqqqqqqqqqqqqqqbahrrgptgylvldeeqqpsqp 113
                      EQQEPRLQYQCQRRCQE-----QQRQHGRGGDLMNPQRGGSGRYEEGEEKQSDNP
                                                                                                                                                                                                                                                                                                         1991-178048/24.
                                               19;
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                                                                                                        918 AA;
                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 556..626 /label= DNA-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "cysteine-rich
                                                         18.5%;
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Pred. No. 3.7;
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                                               Mismatches
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